

<400> 1

```

gccctttctt atgagcatgc ctgtgttggg ttgacagtga gggtaataat gacttgttgg 60
ttgattgtag atatagggct ctcccttgca aggtaattag gctccitaaa ttacctgtaa 120
gattttcttg ccacagcatc cattctgggtt aggcctgggtga tcttctgagt agtgatagat 180
tggttggtgg tgaggtttac aggtgttccc ttctcttact cctgggtgtg gctacaatca 240
ggtggcgtct agagcagcat gggacagggtg ggtaagggga gtcttctcat tatgcagaag 300
tgatcaactt aaatctctgt cagatctacc tttatgtagc ccggcagtcg cgcggattga 360
cggggctcgc ggcgctgggt tcctgggtctc cgggccaggg ca atg ttc cgc acg      414
                                   Met Phe Arg Thr
                                   1

gca gtg atg atg gcg gcc agc ctg gcg ctg acc ggg gct gtg gtg gct      462
Ala Val Met Met Ala Ala Ser Leu Ala Leu Thr Gly Ala Val Val Ala
   5                10                15                20

cac gcc tac tac ctc aaa cac cag ttc tac ccc act gtg gtg tac ctg      510
His Ala Tyr Tyr Leu Lys His Gln Phe Tyr Pro Thr Val Val Tyr Leu
                25                30                35

acc aag tcc agc ccc agc atg gca gtc ctg tac atc cag gcc ttt gtc      558
Thr Lys Ser Ser Pro Ser Met Ala Val Leu Tyr Ile Gln Ala Phe Val
                40                45                50

ctt gtc ttc ctt ctg ggc aag gtg atg ggc aag gtg ttc ttt ggg caa      606
Leu Val Phe Leu Leu Gly Lys Val Met Gly Lys Val Phe Phe Gly Gln
                55                60                65

ctg agg gca gca gag atg gag cac ctt ctg gaa cgt tcc tgg tac gcc      654
Leu Arg Ala Ala Glu Met Glu His Leu Leu Glu Arg Ser Trp Tyr Ala
                70                75                80

gtc aca gag act tgt ctg gcc ttc acc gtt ttt cgg gat gac ttc agc      702
Val Thr Glu Thr Cys Leu Ala Phe Thr Val Phe Arg Asp Asp Phe Ser
   85                90                95                100

ccc cgc ttt gtt gca ctc ttc act ctt ctt ctc ttc ctc aaa tgt ttc      750
Pro Arg Phe Val Ala Leu Phe Thr Leu Leu Leu Phe Leu Lys Cys Phe
                105                110                115

cac tgg ctg gct gag gac cgt gtg gac ttt atg gaa cgc agc ccc aac      798
His Trp Leu Ala Glu Asp Arg Val Asp Phe Met Glu Arg Ser Pro Asn
                120                125                130

atc tcc tgg ctc ttt cac tgc cgc att gtc tct ctt atg ttc ctc ctg      846
Ile Ser Trp Leu Phe His Cys Arg Ile Val Ser Leu Met Phe Leu Leu

```

135	140	145	
ggc atc ctg gac ttc ctc ttc gtc agc cac gcc tat cac agc atc ctg Gly Ile Leu Asp Phe Leu Phe Val Ser His Ala Tyr His Ser Ile Leu 150 155 160			894
acc cgt ggg gcc tct gtg cag ctg gtg ttt gcc ttt gag tat gcc atc Thr Arg Gly Ala Ser Val Gln Leu Val Phe Gly Phe Glu Tyr Ala Ile 165 170 175 180			942
ctg atg acg atg gtg ctc acc atc ttc atc aag tat gtg ctg cac tcc Leu Met Thr Met Val Leu Thr Ile Phe Ile Lys Tyr Val Leu His Ser 185 190 195			990
gtg gac ctc cag agt gag aac ccc tgg gac aac aag gct gtg tac atg Val Asp Leu Gln Ser Glu Asn Pro Trp Asp Asn Lys Ala Val Tyr Met 200 205 210			1038
ctc tac aca gag ctg ttt aca ggc ttc atc aag gtt ctg ctg tac atg Leu Tyr Thr Glu Leu Phe Thr Gly Phe Ile Lys Val Leu Leu Tyr Met 215 220 225			1086
gcc ttc atg acc atc atg atc aag gtg cac acc ttc cca ctc ttt gcc Ala Phe Met Thr Ile Met Ile Lys Val His Thr Phe Pro Leu Phe Ala 230 235 240			1134
atc cgg ccc atg tac ctg gcc atg aga cag ttc aag aaa gct gtg aca Ile Arg Pro Met Tyr Leu Ala Met Arg Gln Phe Lys Lys Ala Val Thr 245 250 255 260			1182
gat gcc atc atg tct cgc cga gcc atc cgc aac atg aac acc ctg tat Asp Ala Ile Met Ser Arg Arg Ala Ile Arg Asn Met Asn Thr Leu Tyr 265 270 275			1230
cca gat gcc acc cca gag gag ctc cag gca atg gac aat gtc tgc atc Pro Asp Ala Thr Pro Glu Glu Leu Gln Ala Met Asp Asn Val Cys Ile 280 285 290			1278
atc tgc cga gaa gag atg gtg act ggt gcc aag aga ctg ccc tgc aac Ile Cys Arg Glu Glu Met Val Thr Gly Ala Lys Arg Leu Pro Cys Asn 295 300 305			1326
cac att ttc cat acc agc tgc ctg cgc tcc tgg ttc cag cgg cag cag His Ile Phe His Thr Ser Cys Leu Arg Ser Trp Phe Gln Arg Gln Gln 310 315 320			1374
acc tgc ccc acc tgc cgt atg gat gtc ctt cgt gca tcg ctg cca gcg Thr Cys Pro Thr Cys Arg Met Asp Val Leu Arg Ala Ser Leu Pro Ala 325 330 335 340			1422
cag tca cca cca ccc ccg gag cct gcg gat cag ggg cca ccc cct gcc			1470

Gln Ser Pro Pro Pro Pro Glu Pro Ala Asp Gln Gly Pro Pro Pro Ala	
345 350 355	
ccc cac ccc cca cca ctc ttg cct cag ccc ccc aac ttc ccc cag ggc	1518
Pro His Pro Pro Pro Leu Leu Pro Gln Pro Pro Asn Phe Pro Gln Gly	
360 365 370	
ctc ctg cct cct ttt cct cca ggc atg ttc cca ctg tgg ccc ccc atg	1566
Leu Leu Pro Pro Phe Pro Pro Gly Met Phe Pro Leu Trp Pro Pro Met	
375 380 385	
ggc ccc ttt cca cct gtc ccg cct ccc ccc agc tca gga gag gct gtg	1614
Gly Pro Phe Pro Pro Val Pro Pro Pro Pro Ser Ser Gly Glu Ala Val	
390 395 400	
gct cct cca tcc acc agt gca gca gcc ctt tct cgg ccc agt gga gca	1662
Ala Pro Pro Ser Thr Ser Ala Ala Ala Leu Ser Arg Pro Ser Gly Ala	
405 410 415 420	
gct aca acc aca gct gct ggc acc agt gct act gct gct tct gcc aca	1710
Ala Thr Thr Thr Ala Ala Gly Thr Ser Ala Thr Ala Ala Ser Ala Thr	
425 430 435	
gca tct ggc cca ggc tct ggc tct gcc cca gag gct ggc cct gcc cct	1758
Ala Ser Gly Pro Gly Ser Gly Ser Ala Pro Glu Ala Gly Pro Ala Pro	
440 445 450	
ggt ttc ccc ttc cct cct ccc tgg atg ggt atg ccc ctg cct cca ccc	1806
Gly Phe Pro Phe Pro Pro Pro Trp Met Gly Met Pro Leu Pro Pro Pro	
455 460 465	
ttt gcc ttc ccc cca atg cct gtg ccc cct gcg ggc ttt gct ggg ctg	1854
Phe Ala Phe Pro Pro Met Pro Val Pro Pro Ala Gly Phe Ala Gly Leu	
470 475 480	
acc cca gag gag cta cga gct ctg gag ggc cat gag cgg cag cac ctg	1902
Thr Pro Glu Glu Leu Arg Ala Leu Glu Gly His Glu Arg Gln His Leu	
485 490 495 500	
gag gcc cgg ctg cag agc ctg cgt aac atc cac aca ctg ctg gac gcc	1950
Glu Ala Arg Leu Gln Ser Leu Arg Asn Ile His Thr Leu Leu Asp Ala	
505 510 515	
gcc atg ctg cag atc aac cag tac ctc acc gtg ctg gcc tcc ttg ggg	1998
Ala Met Leu Gln Ile Asn Gln Tyr Leu Thr Val Leu Ala Ser Leu Gly	
520 525 530	
ccc ccc cgg cct gcc act tca gtc aac tcc act gag ggg act gcc act	2046
Pro Pro Arg Pro Ala Thr Ser Val Asn Ser Thr Glu Gly Thr Ala Thr	
535 540 545	

aca gtt gtt gct gct gcc tcc tcc acc agc atc cct agc tca gag gcc 2094
 Thr Val Val Ala Ala Ala Ser Ser Thr Ser Ile Pro Ser Ser Glu Ala
 550 555 560

acg acc cca acc cca gga gcc tcc cca cca gcc cct gaa atg gaa agg 2142
 Thr Thr Pro Thr Pro Gly Ala Ser Pro Pro Ala Pro Glu Met Glu Arg
 565 570 575 580

cct cca gct cct gag tca gtg ggc aca gag gag atg cct gag gat gga 2190
 Pro Pro Ala Pro Glu Ser Val Gly Thr Glu Glu Met Pro Glu Asp Gly
 585 590 595

gag ccc gat gca gca gag ctc cgc cgg cgc cgc ctg cag aag ctg gag 2238
 Glu Pro Asp Ala Ala Glu Leu Arg Arg Arg Arg Leu Gln Lys Leu Glu
 600 605 610

tct cct gtt gcc cac tga cactgcccc gccagcccc agcctctgct 2286
 Ser Pro Val Ala His
 615

cttttgagca gccctcgctg gaacatgtcc tgccaccaag tgccagctcc ctctctgtct 2346
 gcaccaggga gtagtaccac cagctctgag aaagaggcgg catcccctag gccaaagtga 2406
 aagaggctgg ggttcccatt tgactccagt ccagggcagc catggggatc tcgggtcagt 2466
 tccagccttc ctctccaact cttcagccct gtgtttctgct ggggccaatga aggcagaagg 2526
 tttagcctct gagaagccct cttcttcccc cacccttttc caggagaagg ggctgcccct 2586
 ccaagcccta cttgtatgtg cggagtcaca ctgcagtgcc gaacagtatt agctcccgtt 2646
 cccaagtgtg gactccagag gggctggagg caagctatga acttgctcgc tggcccaccc 2706
 ctaagactgg taccattttc cttttcttac cctgatctcc ccagaagcct cttgtgggtg 2766
 tggctgtgcc ccctatgcc tgtggcattt ctgctctta ctggcaacca cacaactcag 2826
 ggaaaggaat gcctgggagt gggggtgcag gcgggcagca ctgagggacc ctgccccgcc 2886
 cctcccccca ggcccccttc ccctgcagct tctcaagtga gactgacctg tctaccacag 2946
 cagccactgc ccagccgcac tccaggcaag ggccagtgcg cctgctcctg accactgcaa 3006
 tcccagcgcc caaggaaggc cacttctcaa ctggcagaac ttctgaagtt tagaattgga 3066
 attacttcct tactagtgtc ttttggctta aattttgtct ttggaagttg aatgcttaat 3126
 cccgggaaag aggaacagga gtgccagact cctggctctt ccagtttaga aaaggctctg 3186
 tgccaaggag ggaccacagg agctgggacc tgcctgcccc tgtcctttcc ccttggtttt 3246

gtgttacaag agttgttgga gacagtttca gatgattatt taatttgtaa atattgtaca 3306
aattttaata gcttaaattg tatatacagc caaataaaaa ctigcattaa caaaaaaaaa 3366
aaaaaaaa 3374

<210> 2
<211> 617
<212> PRT
<213> Homo sapiens

<400> 2
Met Phe Arg Thr Ala Val Met Met Ala Ala Ser Leu Ala Leu Thr Gly
1 5 10 15
Ala Val Val Ala His Ala Tyr Tyr Leu Lys His Gln Phe Tyr Pro Thr
20 25 30
Val Val Tyr Leu Thr Lys Ser Ser Pro Ser Met Ala Val Leu Tyr Ile
35 40 45
Gln Ala Phe Val Leu Val Phe Leu Leu Gly Lys Val Met Gly Lys Val
50 55 60
Phe Phe Gly Gln Leu Arg Ala Ala Glu Met Glu His Leu Leu Glu Arg
65 70 75 80
Ser Trp Tyr Ala Val Thr Glu Thr Cys Leu Ala Phe Thr Val Phe Arg
85 90 95
Asp Asp Phe Ser Pro Arg Phe Val Ala Leu Phe Thr Leu Leu Leu Phe
100 105 110
Leu Lys Cys Phe His Trp Leu Ala Glu Asp Arg Val Asp Phe Met Glu
115 120 125
Arg Ser Pro Asn Ile Ser Trp Leu Phe His Cys Arg Ile Val Ser Leu
130 135 140
Met Phe Leu Leu Gly Ile Leu Asp Phe Leu Phe Val Ser His Ala Tyr
145 150 155 160
His Ser Ile Leu Thr Arg Gly Ala Ser Val Gln Leu Val Phe Gly Phe
165 170 175
Glu Tyr Ala Ile Leu Met Thr Met Val Leu Thr Ile Phe Ile Lys Tyr
180 185 190
Val Leu His Ser Val Asp Leu Gln Ser Glu Asn Pro Trp Asp Asn Lys

195	200	205
Ala Val Tyr Met Leu Tyr Thr Glu Leu Phe Thr Gly Phe Ile Lys Val		
210	215	220
Leu Leu Tyr Met Ala Phe Met Thr Ile Met Ile Lys Val His Thr Phe		
225	230	235 240
Pro Leu Phe Ala Ile Arg Pro Met Tyr Leu Ala Met Arg Gln Phe Lys		
	245	250 255
Lys Ala Val Thr Asp Ala Ile Met Ser Arg Arg Ala Ile Arg Asn Met		
	260	265 270
Asn Thr Leu Tyr Pro Asp Ala Thr Pro Glu Glu Leu Gln Ala Met Asp		
	275	280 285
Asn Val Cys Ile Ile Cys Arg Glu Glu Met Val Thr Gly Ala Lys Arg		
	290	295 300
Leu Pro Cys Asn His Ile Phe His Thr Ser Cys Leu Arg Ser Trp Phe		
	305	310 315 320
Gln Arg Gln Gln Thr Cys Pro Thr Cys Arg Met Asp Val Leu Arg Ala		
	325	330 335
Ser Leu Pro Ala Gln Ser Pro Pro Pro Pro Glu Pro Ala Asp Gln Gly		
	340	345 350
Pro Pro Pro Ala Pro His Pro Pro Pro Leu Leu Pro Gln Pro Pro Asn		
	355	360 365
Phe Pro Gln Gly Leu Leu Pro Pro Phe Pro Pro Gly Met Phe Pro Leu		
	370	375 380
Trp Pro Pro Met Gly Pro Phe Pro Pro Val Pro Pro Pro Pro Ser Ser		
	385	390 395 400
Gly Glu Ala Val Ala Pro Pro Ser Thr Ser Ala Ala Ala Leu Ser Arg		
	405	410 415
Pro Ser Gly Ala Ala Thr Thr Thr Ala Ala Gly Thr Ser Ala Thr Ala		
	420	425 430
Ala Ser Ala Thr Ala Ser Gly Pro Gly Ser Gly Ser Ala Pro Glu Ala		
	435	440 445
Gly Pro Ala Pro Gly Phe Pro Phe Pro Pro Pro Trp Met Gly Met Pro		
	450	455 460
Leu Pro Pro Pro Phe Ala Phe Pro Pro Met Pro Val Pro Pro Ala Gly		

465				470				475				480				
Phe	Ala	Gly	Leu	Thr	Pro	Glu	Glu	Leu	Arg	Ala	Leu	Glu	Gly	His	Glu	
				485					490					495		
Arg	Gln	His	Leu	Glu	Ala	Arg	Leu	Gln	Ser	Leu	Arg	Asn	Ile	His	Thr	
				500					505					510		
Leu	Leu	Asp	Ala	Ala	Met	Leu	Gln	Ile	Asn	Gln	Tyr	Leu	Thr	Val	Leu	
				515					520					525		
Ala	Ser	Leu	Gly	Pro	Pro	Arg	Pro	Ala	Thr	Ser	Val	Asn	Ser	Thr	Glu	
				530					535					540		
Gly	Thr	Ala	Thr	Thr	Val	Val	Ala	Ala	Ala	Ser	Ser	Thr	Ser	Ile	Pro	
545					550					555					560	
Ser	Ser	Glu	Ala	Thr	Thr	Pro	Thr	Pro	Gly	Ala	Ser	Pro	Pro	Ala	Pro	
				565					570					575		
Glu	Met	Glu	Arg	Pro	Pro	Ala	Pro	Glu	Ser	Val	Gly	Thr	Glu	Glu	Met	
				580					585					590		
Pro	Glu	Asp	Gly	Glu	Pro	Asp	Ala	Ala	Glu	Leu	Arg	Arg	Arg	Arg	Leu	
				595					600					605		
Gln	Lys	Leu	Glu	Ser	Pro	Val	Ala	His								
				610					615							